

## Unidentified Hemoglobin Traits

Each year thousand of infants born in the United States are detected with unidentified hemoglobin variants through newborn screening. Most of these infants are heterozygotes, with the quantity of adult hemoglobin (Hb A) being equal to or greater than the quantity of unidentified hemoglobin variant (Hb U). Mutations in the  $\alpha$ -,  $\beta$ -, or  $\gamma$ -globin genes may cause these variants.

Most unidentified hemoglobin variants have no clinical or hematological consequence, but a few may show altered oxygen affinity or be chemically unstable. Some unidentified hemoglobin variants can cause sickle cell disease when co-inherited with hemoglobin S, but most unidentified hemoglobin variants have no significant genetic implications.

With over 600 structural hemoglobin variants, and limited reference laboratory capacity in the United States, the majority of unidentified hemoglobin variants identified by screening cannot be definitively identified. While the overwhelming number of unidentified hemoglobin variants has no clinical or genetic significance, uncertainty about the identity of variants may lead to frustration and anxiety for families and health care providers.

The algorithm, found on the back of this handout, is intended to provide a common-sense approach that should limit laboratory expense and reserve definitive hemoglobin variant identification in reference labs for situations where definitive identification is needed for specific clinical or genetic concerns in a given family.

Adapted from Sickle cell disease in children and adolescents: Diagnosis, guidelines for comprehensive care, and care paths and protocols for management of acute and chronic complications; Lane, P.A., Buchanan, G.R., Hutter, J.J., Austin, R.F., Brittion, H.A., Rogers, Z.R., et al.



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